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OM nucleic - nucleic search, using sw model

Run on: November 12, 2004, 23:22:45 ; Search time 273 Seconds
(without alignments)
9417.106 Million cell updates/sec

Title: US-09-806-302A-1
Perfect score: 476
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Scoring table: IDENTITY NUC
Gapop 10.0 , Gapext 1.0

Searched: 3625171 seqs, 2700493622 residues

Total number of hits satisfying chosen parameters: 7250342

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications NA:*

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- 18: /cgn2_6/ptodata/1/pubpna/US10F_PUBCOMB.seq:*
- 19: /cgn2_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
- 20: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
- 21: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	476	100.0	476	9	US-09-985-911-5
2	434.2	91.2	497	9	US-09-110-716-30
3	431.6	90.7	517	14	US-10-097-340-186
4	431.6	90.7	517	15	US-10-177-293-279
5	431.6	90.7	517	15	US-10-119-431-26
6	431.6	90.7	517	15	US-10-295-027-503
7	431.6	90.7	733	14	US-10-198-846-10282
8	407.4	85.6	491	9	US-09-967-768A-62
9	295.2	62.0	522	10	US-09-814-353-2203
10	295.2	62.0	522	10	US-09-814-353-8543
11	295.2	62.0	636	10	US-09-814-353-14927
12	293.4	61.6	407	14	US-10-198-846-8737

13	291.8	61.3	499	14	US-10-198-846-129	Sequence 129, Appl
14	280	58.8	368	9	US-09-867-701-6508	Sequence 6508, Ap
15	223.8	47.0	495	9	US-09-956-999-5	Sequence 5, Appli
16	223.8	47.0	495	9	US-09-934-054-4	Sequence 4, Appli
17	223.8	47.0	503	9	US-09-110-716-33	Sequence 33, Appl
18	223.8	47.0	503	9	US-09-934-054-11	Sequence 11, Appl
19	223.8	47.0	503	10	US-09-905-673-27	Sequence 27, Appl
20	223.8	47.0	503	14	US-10-042-945-69	Sequence 69, Appl
21	223.8	47.0	503	15	US-10-157-031-55	Sequence 55, Appl
22	223.8	47.0	503	15	US-10-177-293-277	Sequence 277, Appl
23	223.8	47.0	503	15	US-10-096-319-27	Sequence 27, Appl
24	223.8	47.0	503	15	US-10-393-590-3	Sequence 3, Appli
25	223.8	47.0	503	15	US-10-393-567-3	Sequence 3, Appli
26	223.8	47.0	503	15	US-10-394-087-3	Sequence 3, Appli
27	223.8	47.0	503	17	US-10-283-975A-405	Sequence 405, App
28	223.8	47.0	503	18	US-10-427-217A-17	Sequence 17, Appl
29	223.8	47.0	503	18	US-10-427-217A-18	Sequence 18, Appl
30	223.8	47.0	535	10	US-09-975-502A-1	Sequence 1, Appli
31	223.8	47.0	700	14	US-10-198-846-10860	Sequence 10860, A
32	223.8	47.0	751	14	US-10-198-846-8492	Sequence 8492, Ap
33	223.8	47.0	878	14	US-10-198-846-10961	Sequence 10961, A
34	222.2	46.7	503	9	US-09-825-301-73	Sequence 73, Appl
35	222.2	46.7	503	15	US-10-033-527-73	Sequence 73, Appl
36	217	45.6	871	14	US-10-198-846-1659	Sequence 1659, Ap
37	211.6	44.5	429	10	US-09-905-673-49	Sequence 49, Appl
38	211.6	44.5	429	15	US-10-096-319-49	Sequence 49, Appl
39	210	44.1	429	10	US-09-905-673-46	Sequence 46, Appl
40	210	44.1	429	15	US-10-096-319-46	Sequence 46, Appl
41	208.4	43.8	429	10	US-09-905-673-43	Sequence 43, Appl
42	208.4	43.8	429	10	US-09-905-673-44	Sequence 44, Appl
43	208.4	43.8	429	10	US-09-905-673-45	Sequence 45, Appl
44	208.4	43.8	429	10	US-09-905-673-48	Sequence 48, Appl
45	208.4	43.8	429	15	US-10-096-319-43	Sequence 43, Appl

ALIGNMENTS

RESULT 1
US-09-985-911-5
; Sequence 5, Application US/09985911
; Patent No. US20020151012A1
; GENERAL INFORMATION:
; APPLICANT: NI ET AL.
; TITLE OF INVENTION: HUMAN ENDOMETRIAL SPECIFIC STEROID-BINDING FACTOR I, II AND III
; FILE REFERENCE: PF257D3
; CURRENT APPLICATION NUMBER: US/09/985,911
; CURRENT FILING DATE: 2001-11-06
; PRIOR APPLICATION NUMBER: 09/583,169
; PRIOR FILING DATE: 2000-05-30
; PRIOR APPLICATION NUMBER: 09/263,810
; PRIOR FILING DATE: 1999-03-08
; PRIOR APPLICATION NUMBER: 08/821,451
; PRIOR FILING DATE: 1997-03-21
; PRIOR APPLICATION NUMBER: 60/014,724
; PRIOR FILING DATE: 1996-03-21
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 476
; TYPE: DNA
; ORGANISM: human
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (46)..(330)
; OTHER INFORMATION:
; NAME/KEY: sig_peptide
; LOCATION: (46)..(108)
; OTHER INFORMATION:
; NAME/KEY: mat_peptide
; LOCATION: (109)..(330)
; OTHER INFORMATION:
US-09-985-911-5


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Query Match      90.7%; Score 431.6; DB 14; Length 517;
Best Local Similarity 97.4%; Pred. No. 1.1e-131;
Matches 450; Conservative 0; Mismatches 9; Indels 3; Gaps 1;

QY      6 CTGCCACGACGACTGAACACACAGACAGACGCGCCCTCGCCATGAAGCTGCTGATGTCCT 65
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QY      366 AGACTATGGCCAGAACTCATCTGTGATTGCTAGAAACCACTTTCTTCTTGTC 422
Db      385 AGACTATGGCCAGAACTCATCTGTGATTGCTAGAAACCACTTTCTTCTTGTC 444

QY      423 TTTTATGTGGGAAGCTGCTAGACAACTGTTGAAACCTCAATT 464
Db      445 TTTTATGTGGGAAGCTGCTAGACAACTGTTGAAACCTCAATT 486
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RESULT 4

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US-10-177-293-279
; Sequence 279, Application US/10177293
; Publication No. US20030124128A1
; GENERAL INFORMATION:
; APPLICANT: Lillie, James
; APPLICANT: Glatt, Karen
; APPLICANT: Zhao, Xumei
; APPLICANT: Gannavarpu, Manjula
; APPLICANT: Kamatkar, Shubhangi
; APPLICANT: Mertens, Maureen
; APPLICANT: Myer, Vic
; APPLICANT: Wang, Youzhen
; APPLICANT: Xu, Yongyao
; APPLICANT: Hoersch, Sebastian
; APPLICANT: Monahan, John
; APPLICANT: Meyers, Rachel E.
; APPLICANT: Bast Jr., Robert C.
; APPLICANT: Hortobagyi, Gabriel N.
; APPLICANT: Pusztai, Lajos
; APPLICANT: Meric, Funda
; APPLICANT: Sahin, Aysegul
; APPLICANT: Mills, Gordon B.
; TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF BREAST CANCER
; FILE REFERENCE: MRI-038
; CURRENT APPLICATION NUMBER: US/10/177,293
; CURRENT FILING DATE: 2002-06-21
; PRIOR APPLICATION NUMBER: US 60/299,887
; PRIOR FILING DATE: 2001-06-21
; PRIOR APPLICATION NUMBER: US 60/301,572
; PRIOR FILING DATE: 2001-06-27
; PRIOR APPLICATION NUMBER: US 60/306,501
; PRIOR FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: US 60/325,002
; PRIOR FILING DATE: 2001-09-25
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; PRIOR APPLICATION NUMBER: US 60/362,585
; PRIOR FILING DATE: 2002-03-05
; PRIOR APPLICATION NUMBER: US 60/xxx,xxx
; PRIOR FILING DATE: 2002-05-14
; NUMBER OF SEQ ID NOS: 506
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 279
; LENGTH: 517
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-177-293-279
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Best Local Similarity 97.4%; Pred. No. 1.1e-131;
Matches 450; Conservative 0; Mismatches 9; Indels 3; Gaps 1;

QY      6 CTGCCACGACGACTGAACACACAGACAGACGCGCCCTCGCCATGAAGCTGCTGATGTCCT 65
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QY      66 CATGCTGGCGGCCCTCCTCCTGCACTGCTATGCAGATTCTGGTGCAAACTCCTGGAGGA 125
Db      85 CATGCTGGCGGCCCTCCTCCTGCACTGCTATGCAGATTCTGGTGCAAACTCCTGGAGGA 144

QY      126 CATGTTGAAAAGACCATCAATTCCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 185
Db      145 CATGTTGAAAAGACCATCAATTCCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 204

QY      186 AGAGTTTCATAGACAGTGCCTGCGGCTGAGAGGCTATGGGGAATTCAGGAGTTCCT 245
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QY      366 AGACTATGGCCAGAACTCATCTGTGATTGCTAGAAACCACTTTCTTCTTGTC 422
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QY      423 TTTTATGTGGGAAGCTGCTAGACAACTGTTGAAACCTCAATT 464
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RESULT 5

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US-10-119-431-26
; Sequence 26, Application US/10119431
; Publication No. US20030152939A1
; GENERAL INFORMATION:
; APPLICANT: Smithson, Glennda
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Zhong, Mei
; APPLICANT: Khramtsov, Nikolai
; APPLICANT: Li, Li
; APPLICANT: Gusev, Vladimir
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Anderson, David
; APPLICANT: Shinkets, Richard A.
; TITLE OF INVENTION: NOVEL SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
; FILE REFERENCE: Cura-29 CIP1
; CURRENT APPLICATION NUMBER: US/10/119,431
; CURRENT FILING DATE: 2002-11-15
; PRIOR APPLICATION NUMBER: 60/103,195
; PRIOR FILING DATE: 1998-10-06
; PRIOR APPLICATION NUMBER: 60/282,548
; PRIOR FILING DATE: 2001-04-09
; PRIOR APPLICATION NUMBER: 09/412,231
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; PRIOR FILING DATE: 1999-10-05
; NUMBER OF SEQ ID NOS: 46
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 26
; LENGTH: 517
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-119-431-26

Query Match 90.7%; Score 431.6; DB 15; Length 517;
Best Local Similarity 97.4%; Pred. No. 1.1e-131;
Matches 450; Conservative 0; Mismatches 9; Indels 3; Gaps 1;

QY 6 CTGCCACGCACGACTGAACACACAGACAGCAGCGCCCTCGCCATGAAGCTGCTGATGGTCTT 65
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Db 385 AGACTATGGCCAGAACTCATCTGTTGATTGCTAGAAACCACTTTCTTCTTGTGTTGTC 444
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RESULT 6
US-10-295-027-503
; Sequence 503, Application US/10295027
; Publication No. US20030232350A1
; GENERAL INFORMATION:
; APPLICANT: Afar, Daniel
; APPLICANT: Aziz, Natasha
; APPLICANT: Ginsberg, Wendy M.
; APPLICANT: Gish, Kurt C.
; APPLICANT: Glynn, Richard
; APPLICANT: Hevezi, Peter A.
; APPLICANT: Mack, David H.
; APPLICANT: Murray, Richard
; APPLICANT: Watson, Susan R.
; APPLICANT: Eos Biotechnology, Inc.
; TITLE OF INVENTION: Methods of Diagnosis of Cancer, Compositions and
; TITLE OF INVENTION: Methods of Screening for Modulators of Cancer
; FILE REFERENCE: 018501-012500US
; CURRENT APPLICATION NUMBER: US/10/295,027
; CURRENT FILING DATE: 2002-11-13
; PRIOR APPLICATION NUMBER: US 09/663,733
; PRIOR FILING DATE: 2000-09-15
; PRIOR APPLICATION NUMBER: US 60/350,666
; PRIOR FILING DATE: 2001-11-13
; PRIOR APPLICATION NUMBER: US 60/335,394
; PRIOR FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/332,464

; PRIOR FILING DATE: 2001-11-21
; PRIOR APPLICATION NUMBER: US 60/334,393
; PRIOR FILING DATE: 2001-11-29
; PRIOR APPLICATION NUMBER: US 60/340,376
; PRIOR FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/347,211
; PRIOR FILING DATE: 2002-01-08
; PRIOR APPLICATION NUMBER: US 60/347,349
; PRIOR FILING DATE: 2002-01-10
; PRIOR APPLICATION NUMBER: US 60/355,250
; PRIOR FILING DATE: 2002-02-08
; PRIOR APPLICATION NUMBER: US 60/356,714
; PRIOR FILING DATE: 2002-02-13
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 1386
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 503
; LENGTH: 517
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-295-027-503

Query Match 90.7%; Score 431.6; DB 15; Length 517;
Best Local Similarity 97.4%; Pred. No. 1.1e-131;
Matches 450; Conservative 0; Mismatches 9; Indels 3; Gaps 1;

QY 6 CTGCCACGCACGACTGAACACACAGACAGCAGCGCCCTCGCCATGAAGCTGCTGATGGTCTT 65
Db 25 CTGCCACGCACGACTGAACACACAGACAGCAGCGCCCTCGCCATGAAGCTGCTGATGGTCTT 84
QY 66 CATGCTGGCGCCCTCTCTGCACTGCTATGCGAGATTCTGGCTGCAAACTCTCTGGAGGA 125
Db 85 CATGCTGGCGCCCTCTCTGCACTGCTATGCGAGATTCTGGCTGCAAACTCTCTGGAGGA 144
QY 126 CATGTTGAAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 185
Db 145 CATGTTGAAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 204
QY 186 AGAGTTCATAGACAGTATGCGGCTGCGAGGCTATGGGGAATTCAGCAGTGTTCCT 245
Db 205 AGAGTTCATAGACAGTATGCGGCTGCGAGGCTATGGGGAATTCAGCAGTGTTCCT 264
QY 246 CAACCACTCACATAGAACTCTGAAAAAATTTGGACTGATGATGCATACAGTGTACGACAG 305
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QY 306 CATTTGGTGAATATGAAGAGTAATTAACCTTTACCAAGCGGTTTGGCTCAGAGGGCTAC 365
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QY 366 AGACTATGGCCAGAACTCATCTGTTGATTGCTAGAAACCACTTTCTTCTTGTG---TTGC 422
Db 385 AGACTATGGCCAGAACTCATCTGTTGATTGCTAGAAACCACTTTCTTCTTGTGTTGTC 444
QY 423 TTTTATGTGGAACTGCTAGACAACTGTTGAAACCTCAATT 464
Db 445 TTTTATGTGGAACTGCTAGACAACTGTTGAAACCTCAAAAT 486

RESULT 7
US-10-198-846-10282
; Sequence 10282, Application US/10198846
; Publication No. US20030099974A1
; GENERAL INFORMATION:
; APPLICANT: Lillie, James
; APPLICANT: Xu, Yongyao
; APPLICANT: Wang, Youzhen
; APPLICANT: Steinmann, Kathleen
; TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
; TITLE OF INVENTION: FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
; TITLE OF INVENTION: THERAPY OF BREAST CANCER
; FILE REFERENCE: MRI-049
; CURRENT APPLICATION NUMBER: US/10/198,846

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; CURRENT FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/306,220
; PRIOR FILING DATE: 2001-07-18
; NUMBER OF SEQ ID NOS: 14084
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10282
; LENGTH: 733
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 1, 2, 731, 732, 733
; OTHER INFORMATION: n = A,T,C or G
US-10-198-846-10282

Query Match          90.7%; Score 431.6; DB 14; Length 733;
Best Local Similarity 97.4%; Pred. No. 1.4e-131;
Matches 450; Conservative 0; Mismatches 9; Indels 3; Gaps 1;

QY 6 CTGCCACGACGACTGAACACACAGACAGACGCGCCCTCGCCCATGAAGCTGCTGATGCTCT 65
Db 79 CTGCCACGACGACTGAACACACAGACAGACGCGCCCTCGCCCATGAAGCTGCTGATGCTCT 138
QY 66 CATGCTGGCGGCCCTCTCTGCACTGCTATGCAATTCGAGATTCGGCTGCAAACTCCTGGAGGA 125
Db 139 CATGCTGGCGGCCCTCTCTGCACTGCTATGCAATTCGAGATTCGGCTGCAAACTCCTGGAGGA 198
QY 126 CATGGTTGAAAAGACCATCAATTCGACATATCTATACCTGATGAGGAAATCAAGAGCTTCTTCA 185
Db 199 CATGGTTGAAAAGACCATCAATTCGACATATCTATACCTGATGAGGAAATCAAGAGCTTCTTCA 258
QY 186 AGAGTTCAATAGACAGTGCCTGCGCTGAGAGGCTATGGGAAATCAAGAGCTTCTTCT 245
Db 259 AGAGTTCAATAGACAGTGCCTGCGCTGAGAGGCTATGGGAAATCAAGAGCTTCTTCT 318
QY 246 CAACAGCTCACATAGAACTCTGAAAACCTTTGGACTGATGATGATGATGATGATGATGATGAT 305
Db 319 CAACAGCTCACATAGAACTCTGAAAACCTTTGGACTGATGATGATGATGATGATGATGATGAT 378
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Db 379 CATTGGTGTAAATGAAAGAGTAATTAACCTTTACCCAAAGCGTTGGCTCAGAGGGCTAC 438
QY 366 AGACTATGGCCAGAACTCATCTGTTGATGCTAGAAACCACTTTCTTTGTTG---TTGC 422
Db 439 AGACTATGGCCAGAACTCATCTGTTGATGCTAGAAACCACTTTCTTTGTTGTTGTTGTC 498
QY 423 TTTTATGTGGGAACTGCTAGACAACTGTTGAAACCTCAATT 464
Db 499 TTTTATGTGGGAACTGCTAGACAACTGTTGAAACCTCAAAAT 540

RESULT 8
US-09-967-768A-62
; Sequence 62, Application US/09967768A
; Patent No. US20020150877A1
; GENERAL INFORMATION:
; APPLICANT: Augustus, Meena
; TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
; TITLE OF INVENTION: Sets
; FILE REFERENCE: 689290-72
; CURRENT APPLICATION NUMBER: US/09/967,768A
; PRIOR FILING DATE: 2001-09-28
; PRIOR APPLICATION NUMBER: US/60/236,109
; PRIOR FILING DATE: 2000-09-28
; PRIOR APPLICATION NUMBER: US/60/236,034
; PRIOR FILING DATE: 2000-09-28
; PRIOR APPLICATION NUMBER: US/60/236,111
; PRIOR FILING DATE: 2000-09-28
; NUMBER OF SEQ ID NOS: 325
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 62
; LENGTH: 491
; ORGANISM: Homo sapiens
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-967-768A-62

Query Match          85.6%; Score 407.4; DB 9; Length 491;
Best Local Similarity 97.0%; Pred. No. 1.1e-123;
Matches 426; Conservative 0; Mismatches 11; Indels 2; Gaps 1;

QY 28 GACAGCAGCGCCTCGCCATGAAGCTGCTGATGCTGCTCATGCTGGCGCCCTCCTCCTG 87
Db 1 GACAGCAGCGCCTCGCCATGAAGCTGCTGATGCTGCTCATGATTTGGCGCCCTCCTCCTG 60
QY 88 CACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGACATGGTTGAAAAGACCATCAAT 147
Db 61 CACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGACATGGTTGAAAAGACCATCAAT 120
QY 148 TCCGACATATCTATACCTGAATACAAAGAGCTTCTTCAAGAGTTTCATAGACAGTGATGCC 207
Db 121 TCCGACATATCTATACCTGAATACAAAGAGCTTCTTCAAGAGTTTCATAGACAGTGATGCC 180
QY 208 GCTGCAGAGGCTATGGGAAATTCAGCAGTGTTCCTCAACCACTGTCATAGAACTCTG 267
Db 181 GCTGCAGAGGCTATGGGAAATTCAGCAGTGTTCCTCAACCACTGTCATAGAACTCTG 240
QY 268 AAAAACTTTGGACTGATGATGCATACAGTGTACGACAGCATTTGGTGTAAATATGAAGAGT 327
Db 241 AAAAACTTTGGACTGATGATGCATACAGTGTACGACAGCATTTGGTGTAAATATGAAGAGT 300
QY 328 AATTAACCTTTACCCAGCGCTTTGGCTCAGAGGGCTACAGACTATGGCCAGAACTCATCT 387
Db 301 AATTAACCTTTACCCAGCGCTTTGGCTCAGAGGGCTACAGACTATGGCCAGAACTCATCT 360
QY 388 GTTGATTGCTAGAAACCACTTT--CTTCTTGTGTGTTGTTTATGTGGGAACTGCTAGAC 445
Db 361 GTTGATTGCTAGAAACCACTTTTCTTCTTGTGTGTTGTTTATGTGGGAACTGCTAGAC 420
QY 446 AACTGTTGAAACCTCAATT 464
Db 421 AACTGTTGAAACCTCAAAAT 439

RESULT 9
US-09-814-353-2203/c
; Sequence 2203, Application US/09814353
; Publication No. US20030165831A1
; GENERAL INFORMATION:
; APPLICANT: Lee, John
; APPLICANT: Thompson, Pamela
; APPLICANT: Lillie, James
; TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR
; TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND
; TITLE OF INVENTION: THERAPY OF OVARIAN CANCER
; FILE REFERENCE: MRI-006B
; CURRENT APPLICATION NUMBER: US/09/814,353
; CURRENT FILING DATE: 2001-03-21
; PRIOR APPLICATION NUMBER: US 60/191,031
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: US 60/207,124
; PRIOR FILING DATE: 2000-05-25
; PRIOR APPLICATION NUMBER: US 60/211,940
; PRIOR FILING DATE: 2000-06-15
; PRIOR APPLICATION NUMBER: US 60/216,820
; PRIOR FILING DATE: 2000-07-07
; PRIOR APPLICATION NUMBER: US 60/220,661
; PRIOR FILING DATE: 2000-07-25
; PRIOR APPLICATION NUMBER: US 60/257,672
; PRIOR FILING DATE: 2000-12-21
; NUMBER OF SEQ ID NOS: 22037
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2203
; LENGTH: 522
; TYPE: DNA
; ORGANISM: Homo sapiens
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Db 363 AGAGTTTCATAGACAGTGCCTGCGCTGCAGAGGCTATGGGAAATTCAGCAGTGTTCCT 304
Qy 246 CAACCAAGTCACATAGAACTCTGAAAACTTTGGACTGATGATGCATACAGTGTACGACAG 305
Db 303 CAACCAAGTCACATAGAACTCTGAAAACTTTGGACTGATGATGCATACAGTGTACAGAG 244

RESULT 12
US-10-198-846-8737
; Sequence 8737, Application US/10198846
; Publication No. US2003009974A1
; GENERAL INFORMATION:
; APPLICANT: Lillie, James
; APPLICANT: Xu, Yongyao
; APPLICANT: Wang, Youzhen
; APPLICANT: Steinmann, Kathleen
; TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
; TITLE OF INVENTION: FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
; TITLE OF INVENTION: THERAPY OF BREAST CANCER

; FILE REFERENCE: MRI-049
; CURRENT APPLICATION NUMBER: US/10/198,846
; CURRENT FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/306,220
; PRIOR FILING DATE: 2001-07-18
; NUMBER OF SEQ ID NOS: 14084
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8737
; LENGTH: 407
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 5
; OTHER INFORMATION: n = A,T,C or G
US-10-198-846-8737

Query Match 61.6%; Score 293.4; DB 14; Length 407;
Best Local Similarity 99.7%; Pred. No. 4.8e-86;
Matches 294; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 6 CTGCCACGCACGACTGAACACACAGACAGCAGCCGCTCGCCCATGAAGCTGCTGATGTCCT 65
Db 92 CTGCCACGCACGACTGAACACACAGACAGCAGCCGCTCGCCCATGAAGCTGCTGATGTCCT 151
Qy 66 CATGCTGGCGGCCCTCCTCCTGCACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGGA 125
Db 152 CATGCTGGCGGCCCTCCTCCTGCACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGGA 211
Qy 126 CATGTTGAAAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 185
Db 212 CATGTTGAAAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 271
Qy 186 AGAGTTCATAGACAGTGCCTGCGCTGCAGAGGCTATGGGAAATTCAGCAGTGTTCCT 245
Db 272 AGAGTTCATAGACAGTGCCTGCGCTGCAGAGGCTATGGGAAATTCAGCAGTGTTCCT 331
Qy 246 CAACCAAGTCACATAGAACTCTGAAAACTTTGGACTGATGATGCATACAGTGTAC 300
Db 332 CAACCAAGTCACATAGAACTCTGAAAACTTTGGACTGATGATGCATACAGTGTAC 386

RESULT 13
US-10-198-846-129
; Sequence 129, Application US/10198846
; Publication No. US2003009974A1
; GENERAL INFORMATION:
; APPLICANT: Lillie, James
; APPLICANT: Xu, Yongyao
; APPLICANT: Wang, Youzhen
; APPLICANT: Steinmann, Kathleen
; TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
; TITLE OF INVENTION: FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
; TITLE OF INVENTION: THERAPY OF BREAST CANCER

; FILE REFERENCE: MRI-049
; CURRENT APPLICATION NUMBER: US/10/198,846
; CURRENT FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/306,220
; PRIOR FILING DATE: 2001-07-18
; NUMBER OF SEQ ID NOS: 14084
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 129
; LENGTH: 499
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 9
; OTHER INFORMATION: n = A,T,C or G
US-10-198-846-129

Query Match 61.3%; Score 291.8; DB 14; Length 499;
Best Local Similarity 99.3%; Pred. No. 1.8e-85;
Matches 293; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 6 CTGCCACGCACGACTGAACACACAGACAGCAGCCGCTCGCCCATGAAGCTGCTGATGTCCT 65
Db 78 CTGCCACGCACGACTGAACACACAGACAGCAGCCGCTCGCCCATGAAGCTGCTGATGTCCT 137
Qy 66 CATGCTGGCGGCCCTCCTCCTGCACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGGA 125
Db 138 CATGCTGGCGGCCCTCCTCCTGCACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGGA 197
Qy 126 CATGTTGAAAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 185
Db 198 CATGCTGAAAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCA 257
Qy 186 AGAGTTCATAGACAGTGCCTGCGCTGCAGAGGCTATGGGAAATTCAGCAGTGTTCCT 245
Db 258 AGAGTTCATAGACAGTGCCTGCGCTGCAGAGGCTATGGGAGATTTCAGCAGTGTTCCT 317
Qy 246 CAACCAAGTCACATAGAACTCTGAAAACTTTGGACTGATGATGCATACAGTGTAC 300
Db 318 CAACCAAGTCACATAGAACTCTGAAAACTTTGGACTGATGATGCATACAGTGTAC 372

RESULT 14
US-09-867-701-6508
; Sequence 6508, Application US/09867701
; Patent No. US20020132237A1
; GENERAL INFORMATION:
; APPLICANT: Aglate, Paul A.
; APPLICANT: Jones, Robert
; APPLICANT: Harlocker, Susan L.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.497
; CURRENT APPLICATION NUMBER: US/09/867,701
; CURRENT FILING DATE: 2001-05-29
; NUMBER OF SEQ ID NOS: 10912
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6508
; LENGTH: 368
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-867-701-6508

Query Match 58.8%; Score 280; DB 9; Length 368;
Best Local Similarity 95.8%; Pred. No. 1.2e-81;
Matches 299; Conservative 0; Mismatches 10; Indels 3; Gaps 1;

Qy 156 ATCTATACCTGAATACAAAGAGCTTCTTCAAGAGTTCATAGACAGTATCCGCTGCAGA 215
Db 2 ACCTATACCTGAATACAAAGAGCTTCTTCAAGAGTTCATAGACAGTATCCGCTGCAGA 61
Qy 216 GGCTATGGGAAATTCAGCAGTGTTCCTCAACCAGTCACATAGAACTCTGAAAACTT 275

Db 62 GGCTATGGGAAATTAAGCAGTGTTCCTCAACCAGTCACATAGAACTCTGAAAAACTT 121
Qy 276 TGGACTGATGATGCATACAGTGTACGACAGCATTTGGTGTAAATATGAAGAGTAATTAAC 335
Db 122 TGGACTGATGATGCATACAGTGTACGACAGCATTTGGTGTAAATATGAAGAGTAATTAAC 181
Qy 336 TTACCAAGGCGTTTGGCTCAGAGGCTACAGACTATGGCCAGAACTCATCTGTGATTG 395
Db 182 TTACCAAGGCGTTTGGCTCAGAGGCTACAGACTATGGCCAGAACTCATCTGTGATTG 241
Qy 396 CTAGAAACCACTTTCTCTTGTG---TTGCTTTTATGTGGAACTGCTAGACAACTGTT 452
Db 242 CTAGAAACCACTTTCTCTTGTGTTGTTCTTTTATGTGGAACTGCTAGACAACTGTT 301
Qy 453 GAAACCTCAATT 464
Db 302 GAAACCTCAAT 313

RESULT 15
US-09-956-999-5
; Sequence 5, Application US/09956999
; Patent No. US20020064792A1
; GENERAL INFORMATION:
; APPLICANT: Lincoln, Stephen
; APPLICANT: Klingler, Tod M.
; APPLICANT: Au-Young, Janice
; APPLICANT: Tang, Y. Tom
; APPLICANT: Gould, Richard
; APPLICANT: Akerblom, Ingrid E. J.
; APPLICANT: Seilhamer, Jeffrey
; APPLICANT: Hawkins, Phillip R.
; APPLICANT: Murry, Lynn E.
; APPLICANT: Delegeane, Angelo M.
; APPLICANT: Levine, Wendy B.
; APPLICANT: Hillman, Jennifer L.
; APPLICANT: Goli, Surya K.
; APPLICANT: Altus, Christina M.
; APPLICANT: Bandman, Olga
; APPLICANT: LaBrie, Samuel T.
; APPLICANT: Shah, Purvi
; TITLE OF INVENTION: Database for Storage and Analysis of
; FILE REFERENCE: 6514-069CON
; CURRENT APPLICATION NUMBER: US/09/956,999
; CURRENT FILING DATE: 2001-09-19
; PRIOR APPLICATION NUMBER: 08/282,955
; PRIOR FILING DATE: 1995-07-29
; PRIOR APPLICATION NUMBER: 08/187,530
; PRIOR FILING DATE: 1994-01-27
; PRIOR APPLICATION NUMBER: 08/179,873
; PRIOR FILING DATE: 1994-01-11
; PRIOR APPLICATION NUMBER: 08/100,523
; PRIOR FILING DATE: 1993-08-03
; PRIOR APPLICATION NUMBER: 08/137,951
; PRIOR FILING DATE: 1993-10-14
; PRIOR APPLICATION NUMBER: 07/977,780
; PRIOR FILING DATE: 1992-11-19
; PRIOR APPLICATION NUMBER: 07/916,491
; PRIOR FILING DATE: 1992-07-17
; PRIOR APPLICATION NUMBER: 08/289,822
; PRIOR FILING DATE: 1994-08-12
; PRIOR APPLICATION NUMBER: 08/581,240
; PRIOR FILING DATE: 1995-12-29
; PRIOR APPLICATION NUMBER: 08/744,026
; PRIOR FILING DATE: 1996-11-05
; PRIOR APPLICATION NUMBER: 08/786,999
; PRIOR FILING DATE: 1997-01-23
; PRIOR APPLICATION NUMBER: 08/822,262
; PRIOR FILING DATE: 1997-03-20
; PRIOR APPLICATION NUMBER: 08/951,750
; PRIOR FILING DATE: 1997-10-16
; NUMBER OF SEQ ID NOS: 10

; SOFTWARE: FastSEQ for Windows Version 3.0

; SEQ ID NO 5
; LENGTH: 495
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-956-999-5
Query Match 47.0%; Score 223.8; DB 9; Length 495;
Best Local Similarity 71.4%; Pred. No. 5.4e-63;
Matches 325; Conservative 0; Mismatches 122; Indels 8; Gaps 2;
Qy 7 TGCCACGACGACTGAACACACAGACAGCAGCCGCTCGCCCATGAAGCTGCTGATGCTCCTC 66
Db 7 TGCCACGCGGACTGAACACCGACAGCAGCAGCCTCACCATGAATGCTGATGCTCCTC 66
Qy 67 ATGCTGGGGCCCTCCTCTGCACTGCTATGCAAGTCTGGCTGCAAACTCCTGGAGGAC 126
Db 67 ATGCTGGGGCCCTCCTCCAGCACTGCTACGAGGCTCTGGCTGCCCTTATTGGAGAAT 126
Qy 127 ATGGTTGAAAAAGACCATCAATTCCGACATATCTATACCTGAATACAAAGAGCTTCTTCAA 186
Db 127 GTGATTTCCAAAGACAATCAATCCCAAGTGTCTAAGACTGAATACAAAGAACTTCTTCAA 186
Qy 187 GAGTTCATAGACAGTATGCGCTGACAGAGGCTATGGGGAATTCAGAGCAGTGTTCCTC 246
Db 187 GAGTTCATAGACGACAATGCCACTACAAATGCCATAGATGAATTTGAAGGAATGTTTCTT 246
Qy 247 AACCAGTCAATAGAACTCTGAAAACTTTGGACTGATGATGATGATGATGATGATGATGATG 306
Db 247 AACCAGTCAATAGAACTCTGAGCAATGTTGAGTGTATGCAATTAATATATATGATGATG 306
Qy 307 ATTTGGTGTAAATATGAAGAGTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 366
Db 307 AGTCTTTGTGATTT-----ATTTAACTTTCTGCAAGACCTTTGGCTCACAGACTGCA 360
Qy 367 GACTATGGCCAGAACTCATCTGTTGATTTGCTAGAAAC--CACTTTCTTCTTGTGTTCTT 424
Db 361 GGGTATGCTGAGAAACCAACTACGGATGCTGCAAAACCAACCACTTCTTCTTCTTCTTCTT 420
Qy 425 TTTATGTTGGAACTGCTAGACAACTGTTGAAACCT 459
Db 421 TTTTACTACAAACTACAGACAATTTGTTGAAACCT 455

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Job time : 274 secs

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OM nucleic - nucleic search, using sw model

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Perfect score: 476
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Scoring table: IDENTITY NUC
Gapop 10.0 , Gapext 1.0

Searched: 824507 seqs, 355394441 residues

Total number of hits satisfying chosen parameters: 1649014

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Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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4: /cgn2_6/ptodata/1/ina/6B_COMB.seq: *
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq: *
6: /cgn2_6/ptodata/1/ina/backfiles1.seq: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	476	100.0	476	3	US-08-821-451A-5 X Sequence 5, Appli
2	476	100.0	476	3	US-09-263-810-5 X Sequence 5, Appli
3	476	100.0	476	3	US-09-583-169-5 X Sequence 5, Appli
4	431.6	90.7	517	4	US-09-673-395A-33 Sequence 33, Appli
5	223.8	47.0	495	3	US-08-969-987-5 Sequence 5, Appli
6	223.8	47.0	503	1	US-08-455-896-1 Sequence 1, Appli
7	223.8	47.0	503	2	US-08-933-149-1 Sequence 1, Appli
8	223.8	47.0	503	2	US-09-082-343-1 Sequence 1, Appli
9	223.8	47.0	503	3	US-09-082-253-1 Sequence 1, Appli
10	223.8	47.0	503	4	US-09-162-622-1 Sequence 1, Appli
11	223.8	47.0	503	4	US-09-509-015-1 Sequence 1, Appli
12	223.8	47.0	503	5	PCT-US96-08235-1 Sequence 1, Appli
13	223.8	47.0	535	3	US-09-215-818-1 Sequence 1, Appli
14	223.8	47.0	535	4	US-09-467-602A-1 Sequence 1, Appli
15	201.2	42.3	403	1	US-08-455-896-5 Sequence 5, Appli
16	201.2	42.3	403	2	US-08-933-149-5 Sequence 5, Appli
17	201.2	42.3	403	2	US-09-082-343-5 Sequence 5, Appli
18	201.2	42.3	403	3	US-09-082-253-5 Sequence 5, Appli
19	201.2	42.3	403	4	US-09-162-622-5 Sequence 5, Appli
20	201.2	42.3	403	4	US-09-509-015-5 Sequence 5, Appli
21	201.2	42.3	403	5	PCT-US96-08235-5 Sequence 5, Appli
22	153.4	32.2	279	4	US-09-162-622-15 Sequence 15, Appli
23	153.4	32.2	1233	4	US-09-620-405B-492 Sequence 492, App
24	153.4	32.2	1233	4	US-09-834-759-492 Sequence 492, App
25	153.4	32.2	2232	4	US-09-620-405B-491 Sequence 491, App
26	153.4	32.2	2232	4	US-09-834-759-491 Sequence 491, App
27	153.4	32.2	3288	4	US-09-620-405B-490 Sequence 490, App

28	153.4	32.2	3288	4	US-09-834-759-490	Sequence 490, App
C 29	140.2	29.5	356	4	US-09-389-681-217	Sequence 217, App
C 30	140.2	29.5	356	4	US-09-620-405B-217	Sequence 217, App
C 31	140.2	29.5	356	4	US-09-339-338-217	Sequence 217, App
C 32	140.2	29.5	356	4	US-09-433-826B-217	Sequence 217, App
C 33	140.2	29.5	356	4	US-09-604-287A-217	Sequence 217, App
C 34	140.2	29.5	356	4	US-09-834-759-217	Sequence 217, App
C 35	140.2	29.5	356	4	US-09-590-751A-217	Sequence 217, App
C 36	122.4	25.7	511	4	US-09-389-681-182	Sequence 182, App
C 37	122.4	25.7	511	4	US-09-620-405B-182	Sequence 182, App
C 38	122.4	25.7	511	4	US-09-339-338-182	Sequence 182, App
C 39	122.4	25.7	511	4	US-09-433-826B-182	Sequence 182, App
C 40	122.4	25.7	511	4	US-09-604-287A-182	Sequence 182, App
C 41	122.4	25.7	511	4	US-09-834-759-182	Sequence 182, App
C 42	122.4	25.7	511	4	US-09-590-751A-182	Sequence 182, App
C 43	57.2	12.0	206	1	US-08-455-896-6	Sequence 6, Appli
C 44	57.2	12.0	206	2	US-08-933-149-6	Sequence 6, Appli
C 45	57.2	12.0	206	2	US-09-082-343-6	Sequence 6, Appli

ALIGNMENTS

RESULT 1
US-08-821-451A-5
; Sequence 5, Application US/08821451A
; Patent No. 6066724
; GENERAL INFORMATION:
; APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
; TITLE OF INVENTION: Human Endometrial Specific Steroid-
; TITLE OF INVENTION: Binding Factor I, II and III
; NUMBER OF SEQUENCES: 27
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
; ADDRESSEE: CECCHI, STEWART & OLSTEIN
; STREET: 6 BECKER FARM ROAD
; CITY: ROSELAND
; STATE: NEW JERSEY
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 INCH DISKETTE
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WORD PERFECT 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/821,451A
; FILING DATE: March 21, 1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/014,724
; FILING DATE: March 21, 1996
; ATTORNEY/AGENT INFORMATION:
; NAME: MULLINS, J.G.
; REGISTRATION NUMBER: 33,073
; REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 201-994-1700
; TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 476 BASE PAIRS
; TYPE: NUCLEIC ACID
; STRANDEDNESS: SINGLE
; TOPOLOGY: LINEAR
; MOLECULE TYPE: cdna
US-08-821-451A-5

Query Match 100.0%; Score 476; DB 3; Length 476;
Best Local Similarity 100.0%; Pred. No. 4.6e-145;
Matches 476; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ACAGCTGCCACGACGACTGAACACAGACAGCAGCCCTCGCCATGAAGCTGCTGATG 60

Db	1	ACGAGCTGCCACGACGACTGAACACAGACAGCAGCCGCTCTGCCATGAAGCTGCTGATG	60
Qy	61	GTCCCTCATGCTGGCGGCCCTCCTCCTGCACTGTATGCAGATTCTGGCTGCAAACTCCTG	120
Db	61	GTCCCTCATGCTGGCGGCCCTCCTCCTGCACTGTATGCAGATTCTGGCTGCAAACTCCTG	120
Qy	121	GAGGACATGGTTGAAAAGACCAATCCAATTCGACATATCTATACCTGAATACAAAGAGCTT	180
Db	121	GAGGACATGGTTGAAAAGACCAATCCAATTCGACATATCTATACCTGAATACAAAGAGCTT	180
Qy	181	CTTCAAGAGTTTCATACACAGTGATGCCGCTGCAGAGGCTATGGGGAAATTTCAAGCAGTGT	240
Db	181	CTTCAAGAGTTTCATACACAGTGATGCCGCTGCAGAGGCTATGGGGAAATTTCAAGCAGTGT	240
Qy	241	TTCCCTCAACCAGTCCATAGAACTCTGAAAACTTTGGACCTGATGATGCATACAGTGTTAC	300
Db	241	TTCCCTCAACCAGTCCATAGAACTCTGAAAACTTTGGACCTGATGATGCATACAGTGTTAC	300
Qy	301	GACAGCATTTGGTGTAATATGAAGAGTAATTAACCTTTACCAAGCGCTTTGGCTCAGAGG	360
Db	301	GACAGCATTTGGTGTAATATGAAGAGTAATTAACCTTTACCAAGCGCTTTGGCTCAGAGG	360
Qy	361	GCTACAGACTATGGCCAGAACTCATCTGTTGATTGCTAGAAACCACTTTCTTCTTGTGTT	420
Db	361	GCTACAGACTATGGCCAGAACTCATCTGTTGATTGCTAGAAACCACTTTCTTCTTGTGTT	420
Qy	421	GCTTTTATGTGGGAAGTCTAGACAACTGTTGAAAACCTCAATTCATTCCATTTC	476
Db	421	GCTTTTATGTGGGAAGTCTAGACAACTGTTGAAAACCTCAATTCATTCCATTTC	476

RESULT 2

US-09-263-810-5
; Sequence 5, Application US/09263810
; Patent No. 6174992
; GENERAL INFORMATION:
; APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
; TITLE OF INVENTION: Human Endometrial Specific Steroid-
; TITLE OF INVENTION: Binding Factor I, II and III
; NUMBER OF SEQUENCES: 27
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
; ADDRESSEE: CECCHI, STEWART & OLSTEIN
; STREET: 6 BECKER FARM ROAD
; CITY: ROSELAND
; STATE: NEW JERSEY
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 INCH DISKETTE
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WORD PERFECT 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/263,810
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/821,451
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: MULLINS, J.G.
; REGISTRATION NUMBER: 33,073
; REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 201-994-1700
; TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 476 BASE PAIRS
; TYPE: NUCLEIC ACID
; STRANDEDNESS: SINGLE

; TOPOLOGY: LINEAR		Query Match		100.0%;	Score 476;	DB 3;	Length 476;
; MOLECULE TYPE: cdna		Best Local Similarity		100.0%;	Pred. No. 4.6e-145;		
US-09-263-810-5		Matches 476;		Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;

QY	1	ACGAGCTGCCACGCACGACTGAACACACAGACAGCAGCCCGCCTCGCCATGAAGCTGCTGATG	60
DB	1	ACGAGCTGCCACGCACGACTGAACACACAGACAGCAGCCCGCCTCGCCATGAAGCTGCTGATG	60
QY	61	GTCCCTCATGCTGGCGGCCCTCCTCCTGCACCTGCTATGCAGATTCTGSGTGCAAACTCCTG	120
DB	61	GTCCCTCATGCTGGCGGCCCTCCTCCTGCACCTGCTATGCAGATTCTGSGTGCAAACTCCTG	120
QY	121	GAGGACATGGTTGAAAGACCATCAATTCCGACATATCTATACCTGAATACAAAGAGCTT	180
DB	121	GAGGACATGGTTGAAAGACCATCAATTCCGACATATCTATACCTGAATACAAAGAGCTT	180
QY	181	CTTCAAGAGTTTCATAGACAGTGATGCCGCTGCAGAGGCTATGGGAAATTCAAGCAGTGT	240
DB	181	CTTCAAGAGTTTCATAGACAGTGATGCCGCTGCAGAGGCTATGGGAAATTCAAGCAGTGT	240
QY	241	TTCCCTCAACAGTCACATAGAACTCTGAAAAAATTGGAGCTGATGATGCATACAGTGTAC	300
DB	241	TTCCCTCAACAGTCACATAGAACTCTGAAAAAATTGGAGCTGATGATGCATACAGTGTAC	300
QY	301	GACAGCAATTTGGTGTATATGAAGAGTAATTAACTTTACCCCAAGCGTTTGGCTCAGAGG	360
DB	301	GACAGCAATTTGGTGTATATGAAGAGTAATTAACTTTACCCCAAGCGTTTGGCTCAGAGG	360
QY	361	GCTACAGACTATGGCCGAACCTCATCTGTTGATTGCTAGAAAACCACTTTCTTCTTGTT	420
DB	361	GCTACAGACTATGGCCGAACCTCATCTGTTGATTGCTAGAAAACCACTTTCTTCTTGTT	420
QY	421	GCTTTTATGTGGGAACCTGCTAGACAACCTGTTGAAACCTCAATTTCATTCATTTC	476
DB	421	GCTTTTATGTGGGAACCTGCTAGACAACCTGTTGAAACCTCAATTTCATTCATTTC	476

RESULT 7
US-08-933-149-1
; Sequence 1, Application US/08933149
; Patent No. 5922836
; GENERAL INFORMATION:
; APPLICANT: WATSON, MARK A.
; APPLICANT: FLEMING, TIMOTHY P.
; TITLE OF INVENTION: MAMMAGLOBIN, A SECRETED
; TITLE OF INVENTION: MAMMARY SPECIFIC BREAST CANCER PROTEIN
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS: 14
; ADDRESSEE: HOWELL & HAFFERKAMP, L.C.
; STREET: 7733 FORSYTH BOULEVARD, SUITE 1400
; CITY: ST. LOUIS
; STATE: MISSOURI
; COUNTRY: USA
; ZIP: 63105-1817
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/933,149
; FILING DATE:
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: HENDERSON, MELODIE W.
; REGISTRATION NUMBER: 37,848
; REFERENCE/DOCKET NUMBER: 6029-6040
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (314) 727-5188
; TELEFAX: (314) 727-6092
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 503 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA to mRNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-933-149-1

Query Match 47.0%; Score 223.8; DB 2; Length 503;
Best Local Similarity 71.4%; Pred. No. 6.9e-63;
Matches 325; Conservative 0; Mismatches 122; Indels 8; Gaps 2;
QY 7 TGCCACGCACGACTGAACACACAGACAGCGCGCTCGCCATGAAGCTGCTGATGGTCTC 66
Db 22 TGCCACCGCGACTGAACACCGACGAGCGCTCACCATGAAGTTGCTGATGGTCTC 81
QY 67 ATGCTGGCGGCCCTCTCTGCTGCTATGCGAGATCTGGCTGCAAACTCCTGGAGGAC 126
Db 82 ATGCTGGCGGCCCTCTCCAGCACTGCTACGCGAGCTCTGGCTGCCCTTATTGGAGAAT 141
QY 127 ATGTTGAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCAA 186
Db 142 GTGATTTCCAAGACAATCAATCCCAAGTGTCTAAGACTGAATACAAAGAACTTCTTCAA 201
QY 187 GAGTTCCATAGACGAGTATGCCGCTGCGAGGCTATGGGAAATTCAGGAGCTGTTTCTC 246
Db 202 GAGTTCCATAGACGAGTATGCCGCTGCGAGGCTATGGGAAATTCAGGAGCTGTTTCTC 261
QY 247 AACAGTACATAGAACTCTGAAACTTTTGGACTGATGATGATACAGTGTGACGAGC 306
Db 262 AACCAACCGATGAAGTCTGAGCAATGTTGAGGTGTTTATGCAATTAATATATGACAGC 321
QY 307 ATTTGGTGAATATGAAGAGTAAATTAACCTTTACCAAGCGGTTTGGCTCAGAGGCTACA 366
Db 322 AGCTTTGTGATTT-----ATTTTAACTTTCTGCAAGACCTTTGGCTCAGAGAACTGCA 375

QY 367 GACTATGCCAGAACTCATCTGTTGATGCTAGAAAC--CACTTTCTTCTTGTGTTGCTT 424
Db 376 GGGTATGGTGAGAAACCAACTACGGATGCTGCAAAACCAACACCTTCTTCTTATGTCT 435
QY 425 TTTATGTGGAACTGCTAGACAACTGTTGAAACCT 459
Db 436 TTTTACTACAACTACAAGACAATTGTTGAAACCT 470
RESULT 8
US-09-082-343-1
; Sequence 1, Application US/09082343
; Patent No. 5968754
; GENERAL INFORMATION:
; APPLICANT: WATSON, MARK A.
; APPLICANT: FLEMING, TIMOTHY P.
; TITLE OF INVENTION: DNA SEQUENCE AND ENCODED
; TITLE OF INVENTION: MAMMARY-SPECIFIC BREAST CANCER PROTEIN
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ROGERS, HOWELL & HAFFERKAMP
; STREET: 7733 FORSYTH BOULEVARD, SUITE 1400
; CITY: ST. LOUIS
; STATE: MISSOURI
; COUNTRY: USA
; ZIP: 63105-1817
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/082,343
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/455,896
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: HOLLAND, DONALD R.
; REGISTRATION NUMBER: 35,197
; REFERENCE/DOCKET NUMBER: 952726
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (314) 727-5188
; TELEFAX: (314) 727-6092
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 503 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA to mRNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-09-082-343-1
Query Match 47.0%; Score 223.8; DB 2; Length 503;
Best Local Similarity 71.4%; Pred. No. 6.9e-63;
Matches 325; Conservative 0; Mismatches 122; Indels 8; Gaps 2;
QY 7 TGCCACGCACGACTGAACACACAGACAGCGCGCTCGCCATGAAGCTGCTGATGGTCTC 66
Db 22 TGCCACCGCGACTGAACACCGACGAGCGCTCACCATGAAGTTGCTGATGGTCTC 81
QY 67 ATGCTGGCGGCCCTCTCTGCTGCTGCTATGCGAGTCTTGGCTGCAAACTCCTGGAGGAC 126
Db 82 ATGCTGGCGGCCCTCTCCAGCACTGCTACGCGAGCTCTGGCTGCCCTTATTGGAGAAT 141
QY 127 ATGTTGAAAGACCATCAATTCGACATATCTATACCTGAATACAAAGAGCTTCTTCAA 186
Db 142 GTGATTTCCAAGACAATCAATCCCAAGTGTCTAAGACTGAATACAAAGAACTTCTTCAA 201
QY 187 GAGTTCCATAGACGAGTATGCCGCTGCGAGGCTATGGGAAATTCAGGAGCTGTTTCTC 246

Db 103 ATGCTGGCGGCTCTCCAGCACTGTCTACGAGGCTCTGGCTGCCCTTATTGGAGAAT 162
QY 127 ATGGTTGAAAAGACCATCAATCCGACATATCTATACCTGAATACAAAGAGCTTCTTCAA 186
Db 163 GTGATTTCCAAAGACAATCAATCCACAGTGTCTAAGACTGAATACAAAGAACTTCTTCAA 222
QY 187 GAGTTCATAGACAGTGTGCGCTGCAGAGGCTATGGGAAAATTCAGCAGTGTTCCTC 246
Db 223 GAGTTCATAGACGACAATGCCACTACAAATGCCATAGATGAATGAAGGAATGTTTTCTT 282
QY 247 AACCAGTCACATAGAACTCTGAAAAAATTTGGACTGATGATGCATACAGTGTACGACAGC 306
Db 283 AACCAACCGATGAAACTCTGAGCAATGTGAGGTGTTTATGCAATTAATATATGACAGC 342
QY 307 ATTGGTGAATATGAAGAGTAATTAACCTTACCAGGCGTTTGGCTCAGAGGCTACA 366
Db 343 AGTCTTTGTGATTT-----ATTTTAACTTTCTGCAAGACCTTTGGCTCACAGAACTGCA 396
QY 367 GACTATGCCAGAACTCATCTGTGTTGATGCTAGTCTAGAAAC--CACTTTCTTCTTGTGCTT 424
Db 397 GGGTATGGTGAGAAACAGCTACGGATGCTGCAAAACCACACCTTCTCTTCTTATGTCT 456
QY 425 TTTATGTGGGAACTGCTAGACAACTGTGAAAACCT 459
Db 457 TTTTACTACAACTACAGACAATTTGTGAAAACCT 491

RESULT 15
US-08-455-896-5
; Sequence 5, Application US/08455896
; Patent No. 5668267
; GENERAL INFORMATION:
; APPLICANT: WATSON, MARK A.
; APPLICANT: FLEMING, TIMOTHY P.
; TITLE OF INVENTION: DNA SEQUENCE AND ENCODED
; TITLE OF INVENTION: MAMMARY-SPECIFIC BREAST CANCER PROTEIN
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ROGERS, HOWELL & HAERKAMP
; STREET: 7733 FORSYTH BOULEVARD, SUITE 1400
; CITY: ST. LOUIS
; STATE: MISSOURI
; COUNTRY: USA
; ZIP: 63105-1817
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/455,896
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: HOLLAND, DONALD R.
; REGISTRATION NUMBER: 35,197
; REFERENCE/DOCKET NUMBER: 952726
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (314) 727-5188
; TELEFAX: (314) 727-6092
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 403 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA to mRNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-455-896-5

Query Match 42.3%; Score 201.2; DB 1; Length 403;
Best Local Similarity 71.9%; Pred. No. 1.4e-55;

Matches 279; Conservative 0; Mismatches 103; Indels 6; Gaps 1;
QY 7 TGCCACGCAGCTGAACACAGACAGCAGCGCCCTCGCCATGAAGCTGCTGATGGTCTC 66
Db 22 TGCCACCCGCGACTGAACACCGACAGCAGCGCCTCACCATGAAGTTGCTGATGGTCTC 81
QY 67 ATGCTGGCGGCTCCTCCTGCACTGCTATGCAGATTCTGGCTGCAAACTCCTGGAGGAC 126
Db 82 ATGCTGGCGGCTCCTCCTGCACTGCTATGCAGGCTCTGGCTGCCCTTATTGGAGAAT 141
QY 127 ATGGTTGAAAAGACCATCAATCCGACATATCTATACCTGAATACAAAGAGCTTCTTCAA 186
Db 142 GTGATTTCCAAAGACAATCAATCCACAGTGTCTAAGACTGAATACAAAGAACTTCTTCAA 201
QY 187 GAGTTCATAGACAGTGTGCGCTGCAGAGGCTATGGGAAAATTCAGCAGTGTTCCTC 246
Db 202 GAGTTCATAGACGACAATGCCACTACAAATGCCATAGATGAATGAAGGAATGTTTTCTT 261
QY 247 AACCAGTCACATAGAACTCTGAAAAAATTTGGACTGATGATGCATACAGTGTACGACAGC 306
Db 262 AACCAACCGATGAACTCTGAGCAATGTGAGGTGTTTATGCAATTAATATATGACAGC 321
QY 307 ATTTGGTGAATATGAAGAGTAATTAACCTTACCCAGGCGTTTGGCTCAGAGGCTACA 366
Db 322 AGTCTTTGTGATTT-----ATTTTAACTTTCTGCAAGACCTTTGGCTCACAGAACTGCA 375
QY 367 GACTATGCCAGAACTCATCTGTGTTGATT 394
Db 376 GGGTATGGTGAGAAACCAACTACGGATT 403

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Job time : 60 secs

